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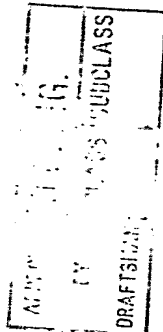
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WO 98/41234

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FIGURE 1(a). [SEQ ID NO:1]

5 ATGACTTTTAAACAATAAACTATTGAAGAGTTGCACAATCTCCTTGTCTCTAAGGAAATT
TCTGCAACAGAAATTGACCCCAAGCAACACTTGAAATAATCAAGTCTCGTGAGGAAGCCATC
AATTCAATTGTCAACCATCGCTGAGGAGCAAGCTCTTGTTCAGGAATTCACCTTGTCTTAAGGATAACATC
GCTGGAATTGATGCTGACAAATGTCCTTTCAGGAATTCACCTTGTCTTAAGGATAACATC
TCTACAGACGGTATTCTCACAACTGCTGCCCTCAAAAATGCTCTACAACTATGAGCCAAATC
TTTGATGCGACAGCTGTGCGCAATGCAAAAACCAAGGCATGATTGTGCTTGGAAAGACC
AACATGGACGAATTTGCTATGGGTGGTTTCAGGTGAAACTTCACACTACGGAGCAACTAAA
AACGCTTGGGACACAGCAAGGTTCCCTGGTGGTCTCAAGTGGTTCATCGGCCCAACCT
GCCTCAGGACAAAGTTGCTTGTCACTTGGTCTCAAAACCAACTACGGAAACAGTTTTCAGTTTCGGT
GCTGCTTCAACGGAATCGTTGGTCTCAAAACCAACTACGGAAACAGTTTTCAGTTTTCGGT
CTCATTTGCCCTTTGGTAGCTCATTAGACCAAGATTGGACCTTTTGTCTCTACTGTTAAGGAA
AATGCCCTCTTGTCAACGCTATTGCCAGCGAAGATGCTAAAGACTCTACTTCTGCTCCT
GTCCGCATCGCCGACTTTACTTCAAAAATCGGCCAAGACATCAAGGATGAAAAATCGCT
TTGCCCTAAGGAATACCTCGGTGAAGGAATTAACCCAGAGGTTAAGGAAACCATTTCTAAAT
15 GCCGCTAAACACTTTGAAAATTTGGTGTCTATTGTCGAAGAAGTCAGCCTTCCCTCACTCT
AAATACGGAGTTGCCGTATACATCATCGCTTCATCAGAAAGCTTCATCAAACTTGCAA
CGCTTCGACGGTATCCGTTACGGCTATCGCGCAGAGATGCAACCACTTGATGAAATC
TATGTAAACAGCCGAAGCCAAAGTTTGGTGAAGAAGTGAAGCGCGCTATCATGCTGGGT
ACTTTCAGTCTTTTCATCAGGTTACTACGATGCCCTACTATAAGAAGCTGGACAGGTCCGT
ACACTTATCATCAAGATTTGCAAAAAGTCTTCGCGGATTACGATTGATTTTGGGTCCA
20 ACTGCTCCAAGTGTGCCCTATGACTTGGATTCTCTCAACCATGACCCAGTTGCCATGTAC
TTAGCCGACCTATTGACCAATACCTGTAAACCTTGGCAGGACTGCCCTGGAAATTCGATTCCCT
GCTGGATTCTCTCAAGGTCTACCTGTGCGACTCCAATTGATTGGTCCCAAGTACTCTGAG
GAAACCATTTACCAAGCTGCTGTGCTTTTGAAGCAACAACAGACTACCAACAACAACAA
25 CCCGTGATTTTGGAGGTGACAAC



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FIGURE 1(b). [SEQ ID NO:3]

TTTGCAACAGAAATTGACCCCAAGCAACACTTGAAATATCAAGTCTCGTGAGGAAGCCATCAATTCATTG

FIGURE 2(a). [SEQ ID NO:2]

MTFNNKTI EELHNLVSK EISATELTQATLENIKSREEAINSFVTIAEEQALVQAKAIDE
 AGIDADNVLSGIPLAVKDNISTDGI LTTAASKMLYNYEPIFDATAVANAKTKGMI VVGKT
 NMDEFAMGSGGETSHYGATKNAWDH SKVPGSSSGSAAA VASGVRLSLGSDTGG SIRQP
 AAFNGIVGLKPTYGTVSRFGLIAFGSSLDQIGPFAPTVKENALLN AIASEDAKDSTSAP
 5 VRIADFTSKIGQDIKGMKIALPK EYLGEGINPEVKETILNAAKHFEKLGAI VEEVSLPHS
 KYGVAVYYIIASSEASSNLQRF DGIRYGYRAEDATNLDEIYVNSRSQGFGE EVKRRIMLG
 TFSLSGGYDAYYKKAGQVRTLI IQDFEKFADYDLILGPTAPSVAYDLD SLNHDPVAMY
 LADLLTIPVNLAGLPGISIPAGFSQGLPVGLQLIGPKYSEETIYQAAAAFEATTDYHKQQ
 PVIFGGDN



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FIGURE 2(b). [SEQ ID NO:4]

FATELTQATLENKSREEAINSFVTIAEEQALVQAKAIDEAGIDADNVLSGIPLAVKDNISTDGILTTAA